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FIG. 1

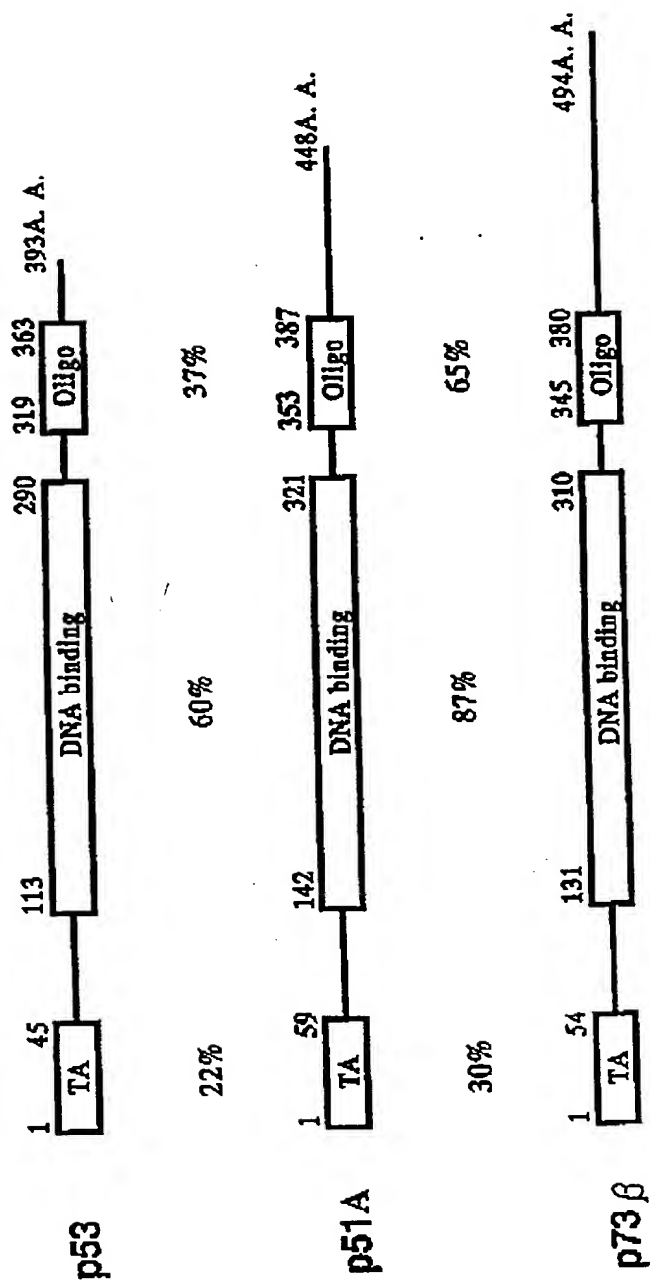


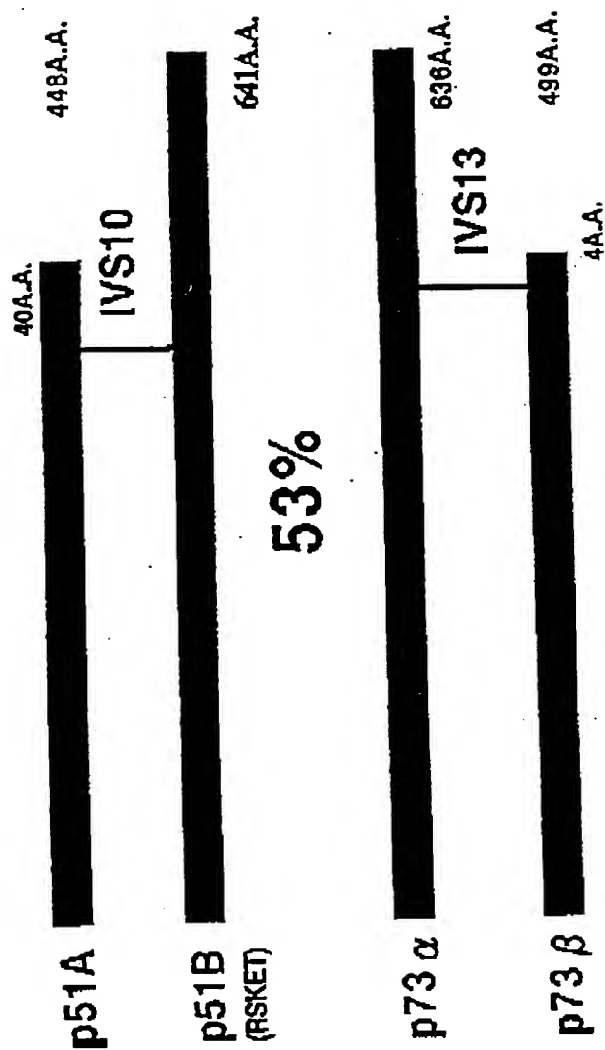
FIG. 2

p51A	MSQSTQTNEF	LSPEVFQHW	DLLEQPTCSV	QPIQUNFVDE	PSDQATNKEI	50
p73B	MAQSTATSP	DGGTTFEHLW	SSLE-POSTY	--FOUP-QSS	RGHNEVVGGT	45
p53	MEEPQSDPS	VEPPLSQETF	SLU-----	--WLL-----L	PENNVLSPLP	36
Consensus	..QST.T...	..P..FQH.W	SLE-P....	---LL.....	P.MN.....	50
p51A	EISMCIHQ	DSLSDPHMP	QYTNLGLLNS	MDQIQONGSS	STSPYNTDHA	100
p73B	DSIMCVFHE	--GHTTSVMA	CF---NLLES	THDQSSRAA	SASPYTPEHA	90
p53	SOAMDLMLS	---POOI-E	ON---FTED	PGPDEAPRMP	EAPF-RYARA	76
Consensus	..SM...L.D....	Q....LL.S	...Q...R..	SASPY...A	100
p51A	QNSVTAPSPY	AGSSSTFDAL	SPSPALPSNT	DYFQPHSDV	IFQSSSTAKS	150
p73B	--ASVPTNSPY	AGSSSTFDTH	SPAPVPSNT	DYFQPHREV	IFQSSSTAKS	139
p53	--PAAPTPAA	PAPAPSW-PL	SSS--VPCK	TYGQSYGRL	GLHSSTAKS	121
Consensus	..SVPTSPSY	AGSSSTFD..L	SPSP..PSNT	DYFQPH..V	IFQSSSTAKS	150
p51A	ATITYSTELK	RLYCCQAKTC	FCQINVTTP	FCQAVIRAMP	VYKAEHTE	200
p73B	ATITYSPULK	RLYCCQAKTC	FCQINVTTP	FCQAVIRAMP	VYKAEHTE	189
p53	VCTITYSPALN	RLYCCQAKTC	FCQINVTTP	FCQAVIRAMP	VYKAEHTE	171
Consensus	ATITYSP..K	RLYCCQAKTC	FCQINVTTP	FCQAVIRAMP	VYKAEHTE	200
p51A	VVRCQPHL	SREFNEQIA	PSHLIRVEG	NSHAGVEEP	TVRCQSWVF	250
p73B	VVRCQPHL	GRDFNEQIA	PSHLIRVEG	NHLSQVQOP	TVRCQSWVF	239
p53	VVRCQPHL	CSD-SQC-IA	PSHLIRVEG	NHLYEYDOR	TVRCQSWVF	219
Consensus	..VRCQPHL	..RDFNEQIA	PSHLIRVEG	..NSHAGVEEP	TVRCQSWVF	250
p51A	YEPPVQTEF	TTILYHFMCH	SSCGGNRR	PILLITLBT	RQCNLGRS	300
p73B	YEPPVQTEF	TTILYHFMCH	SSCGGNRR	PILLITLBT	RQCNLGRS	289
p53	YEPPVQSDC	TTILYHFMCH	SSCGGNRR	PILLITLBT	RQCNLGRS	269
Consensus	YEPPVQTEF	TTILYHFMCH	SSCGGNRR	PILLITLBT	RQCNLGRS	300
p51A	FEARICACPG	RDRKAEDESI	RQQQVSD--S	TKMGQGYKRP	FRQNTHTGI-Q	347
p73B	FEARICACPG	RDRKAEDEHY	REQQALNESS	AKNGAASKRA	FRQSPPAVPA	339
p53	FEARICACPG	RDRRTTEENL	RCKCEPH--H	ELPPGSKRA	LPNHTSS---	314
Consensus	FEARICACPG	RDRKAEDE..	RQQQ....-S	.KNG..YKRA	F.QNT.....	350
p51A	M-TSINKRRS	RDDELLYPY	RGRRTBMLL	KINGELELMQ	YLPQNTIETY	396
p73B	LGAGVKKRH	QEDTYLUQV	RGRRTBMLL	KINGELELMQ	YLPQNTIETY	389
p53	--SPQYKKP	LDGEYFLQI	RGRRTBMLL	ELNEALELKO	AQAGKEPGGS	362
ConsensusNKRR.	..E...YLUQV	RGRRTBMLL	KINGELELMQ	..PQ.....Y	400
p51A	RQQQQQHQ	HL-----	---LQ-H-	---LLSACFRNE	LV-EPRRE--	427
p73B	RQQQQLLRP	HLPPSYGP	VLSPIQVHG	GHNKLPVNO	LVGQPPHSS	439
p53	RAHSS----	HL-----	---K-----	---SKKGQ	---STSRH--	380
Consensus	RQQQQ....	HL-----	---LQ-H-	---NQ	LV-.P.RH--	450
p51A	--TPKQSDV	--FFR--HS	KPPN-----	-----RSVYP-	448	
p73B	AATPHLGPGV	PGMLNHGHA	VPANGESSS	MSAQSMVSGS	HCTPPPPYKA	489
p53	--KKL----	--HFK--T	EGPD-----	---SD-----	393	
Consensus	--TPKL..V	--MF....H	PPN-----	-----S.Y--	500	
p51A	-----	-----	-----	-----	448	
p73B	DPQLVRTWGP	-----	-----	-----	499	
p53	-----	-----	-----	-----	393	
Consensus	-----	-----	-----	-----	510	

FIG. 3

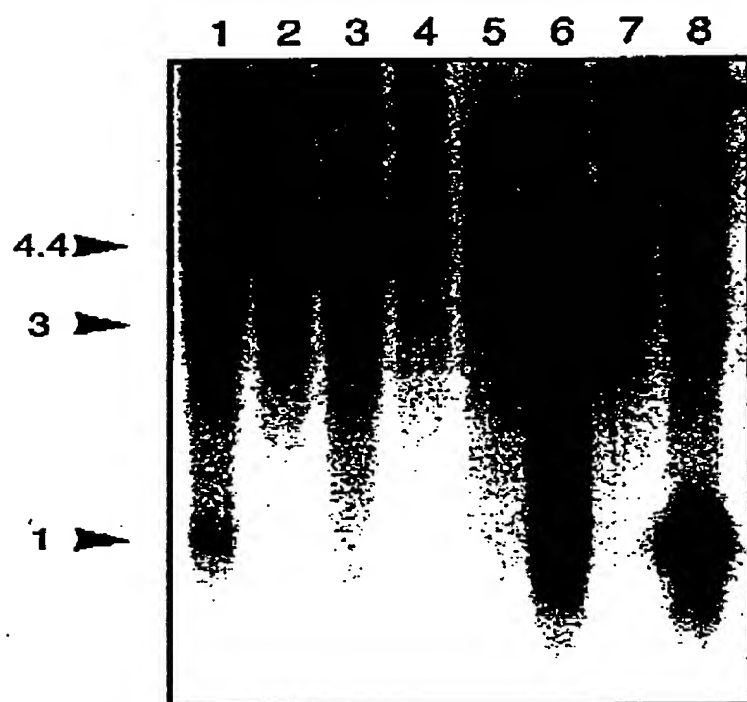
p518	EQSTSTNEF LSPENFCHIR DLEQITCSV QPDKHFVDL PSEDGATNKI	50
p73d	EQSTSTASP DGGTTEHHR ESLEIFOSTY --RQLP-QSS RGNNEVVGGT	45
Consensus	EQSTST... ..ESLEIFOSTY... ..RQLP-QSS	50
p518	E SMCIRMQ DSOLSDPMP QYTHLGLHS MDQIQNGSS S SPTNTQHA	100
p73d	DSMCQVHRE --GHTYSMA QF --ALLSS MDQIQSSRAA SASPTTPHA	90
Consensus	SMC... ..Q... ..L... ..S... ..S SPT... ..HA	100
p518	QSVTAISPY AQPSSTFQAL SPSAIPSKT DYPGPHFQV FQQSSTAKS	150
p73d	ASVPTHSPY AQPSSTFQTM SPSAIPSKT DYPGPHFQV FQQSSTAKS	139
Consensus	SV... ..SPY AQPSSTF... ..SPSAIPSKT DYPGPHFQV FQQSSTAKS	150
p518	ATWTYSTLK KLYCQIARTC PIQIKVTPP PQDANIRAMP VYKKAEMTE	200
p73d	ATWTYSTLK KLYCQIARTC PIQIKVTPP PQDANIRAMP VYKKAEMTD	189
Consensus	ATWTYSTLK KLYCQIARTC PIQIKVTPP PQDANIRAMP VYKKAEMTD	200
p518	VVKRCPMHEL SRFNEGGA FASHLIRVEG NSMAYVEQP VTGRQSVVVF	250
p73d	VVKRCPMHEL SRFNEGGA FASHLIRVEG NMLQYVQDP VTGRQSVVVF	239
Consensus	VVKRCPMHEL SRFNEGGA FASHLIRVEG N... ..QYVQDP VTGRQSVVVF	250
p518	YEPPQVGTEF TILYMFNCH SSCVGGHNR PILIITLET RDGQVLGRRC	300
p73d	YEPPQVGTEF TILYMFNCH SSCVGGHNR PILIITLET RDGQVLGRRC	289
Consensus	YEPPQVGTEF TILYMFNCH SSCVGGHNR PILIITLET RDGQVLGRRC	300
p518	FEARICACPG RDRKADESI RQCVSO--S YKMDGCKRP FQNTHTCI-Q	347
p73d	FEARICACPG RDRKADEHY RQCVSO--S YKMDGCKRP FQNTHTCI-Q	339
Consensus	FEARICACPG RDRKADE... ..RQCVSO--S YKMDGCKRP FQNTHTCI-Q	350
p518	M-TSIKKRP SDELLYLV RGRBTGLL RKESLELMQ YLPQHTIEY	396
p73d	LGAGYKKRP SDELLYLV RGRBTGLL RKESLELMQ YLPQHTIEY	389
ConsensusIKKRP... ..YLV RGRBTGLL RKESLELMQ YLPQHTIEY	400
p518	RQQQQQCHN LLOKQTSICS FSYQNSRR LHHN-SMKN LPSVQINP	445
p73d	RQQQQQCHN LLOKQTSICS FSYQNSRR LHHN-SMKN LPSVQINP	433
Consensus	R... ..Q... ..LLOKQTSICS FSYQNSRR LHHN-SMKN LPSVQINP	450
p518	QQRNALTHT IPOGQANIP LMGTHMPHAG DNGLSPTQA LPPPISTST	495
p73d	PPPHSSAATP RLGPVQPG--MLNNGHAPV ANGENSESH--SAGSHVSG	478
ConsensusHT... ..IP... ..L... ..H... ..S... ..S... ..S	500
p518	SHCTPPPPY TQSTVFLA ELGSSGLQY FQGGUTTY DIENYSMDL	545
p73d	SHCTPPPPY TQSTVFLA ELGSSGLQY FQGGUTTY DIENYSMDL	528
Consensus	SHCTPPPPY TQSTVFLA ELGSSGLQY FQGGUTTY DIENYSMDL	550
p518	ASLKIPEDR MATRQILQH RQHEFSPS HLLRTPSSAS TVSGSSETR	595
p73d	GALKIPECR MATRQILQH RQHEFSPS HLLRTPSSAS TVSGSSETR	578
ConsensusLKIPEDR... ..MATRQILQH RQHEFSPS HLLRTPSSAS TVSGSSETR	600
p518	GRRVIGAVF YRRTTSRP R-----DE RHPHFQMDA RUKKCHKE	638
p73d	RGRVIGAVF YRRTTSRP R-----DE RHPHFQMDA RUKKCHKE	628
ConsensusVIGAVF YRRTTSRP R-----DE RHPHFQMDA RUKKCHKE	650
p518	--RGR--	641
p73d	EFTGRH	636
ConsensusRGR... ..EFTGRH	650

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FIG. 5



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FIG. 6

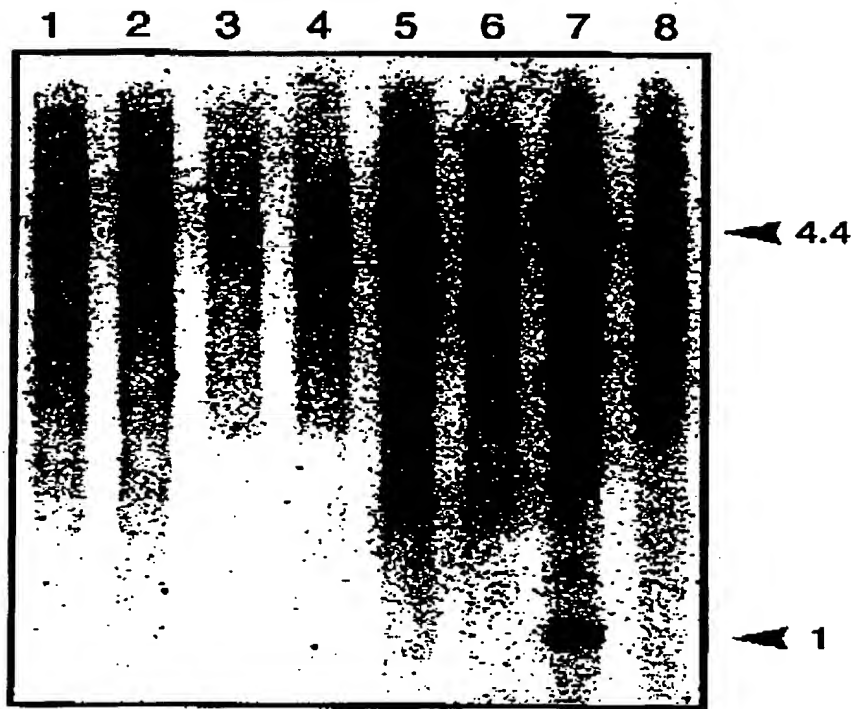


FIG. 7

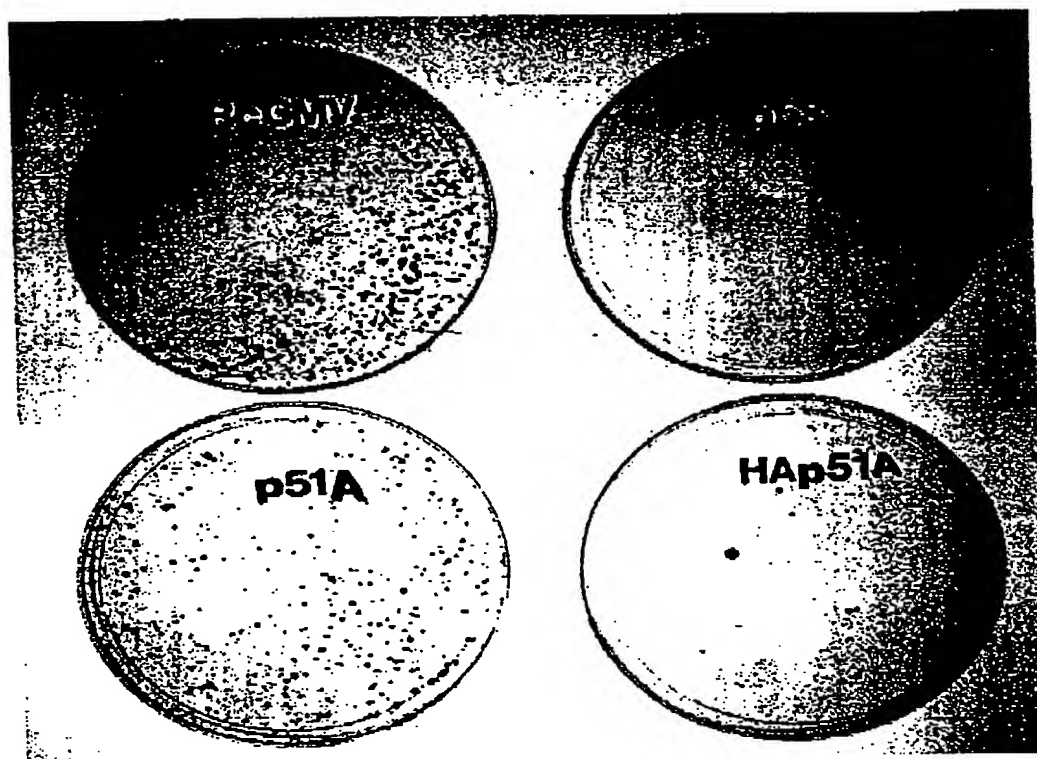


FIG. 8

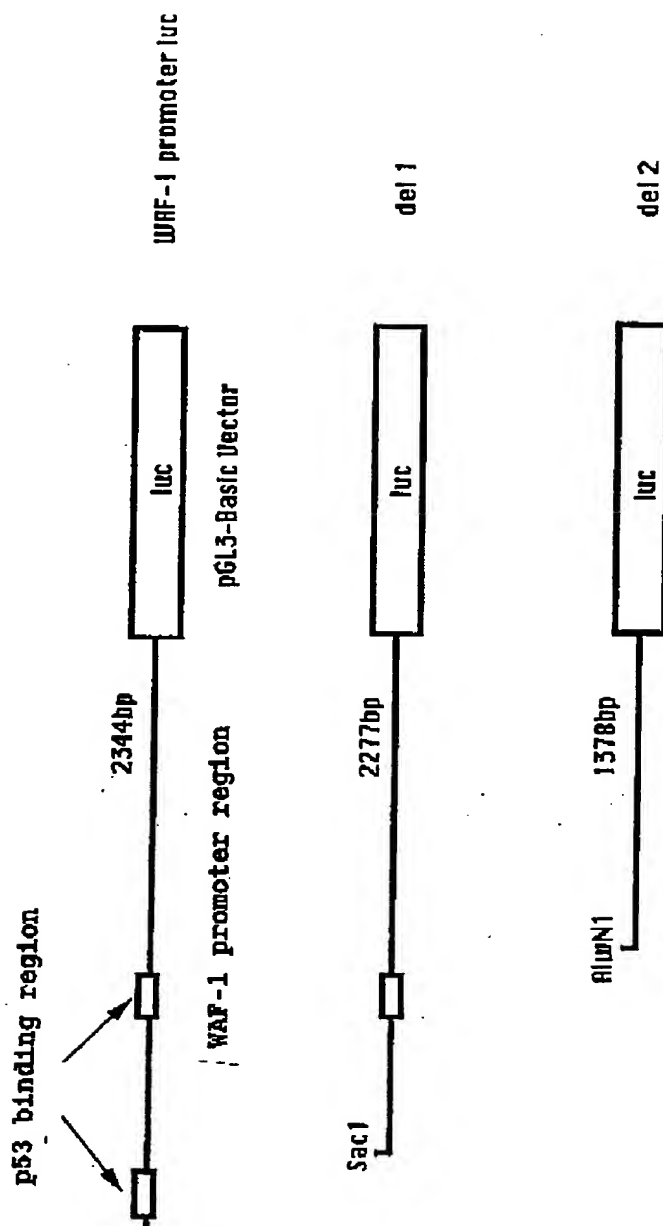


FIG. 9

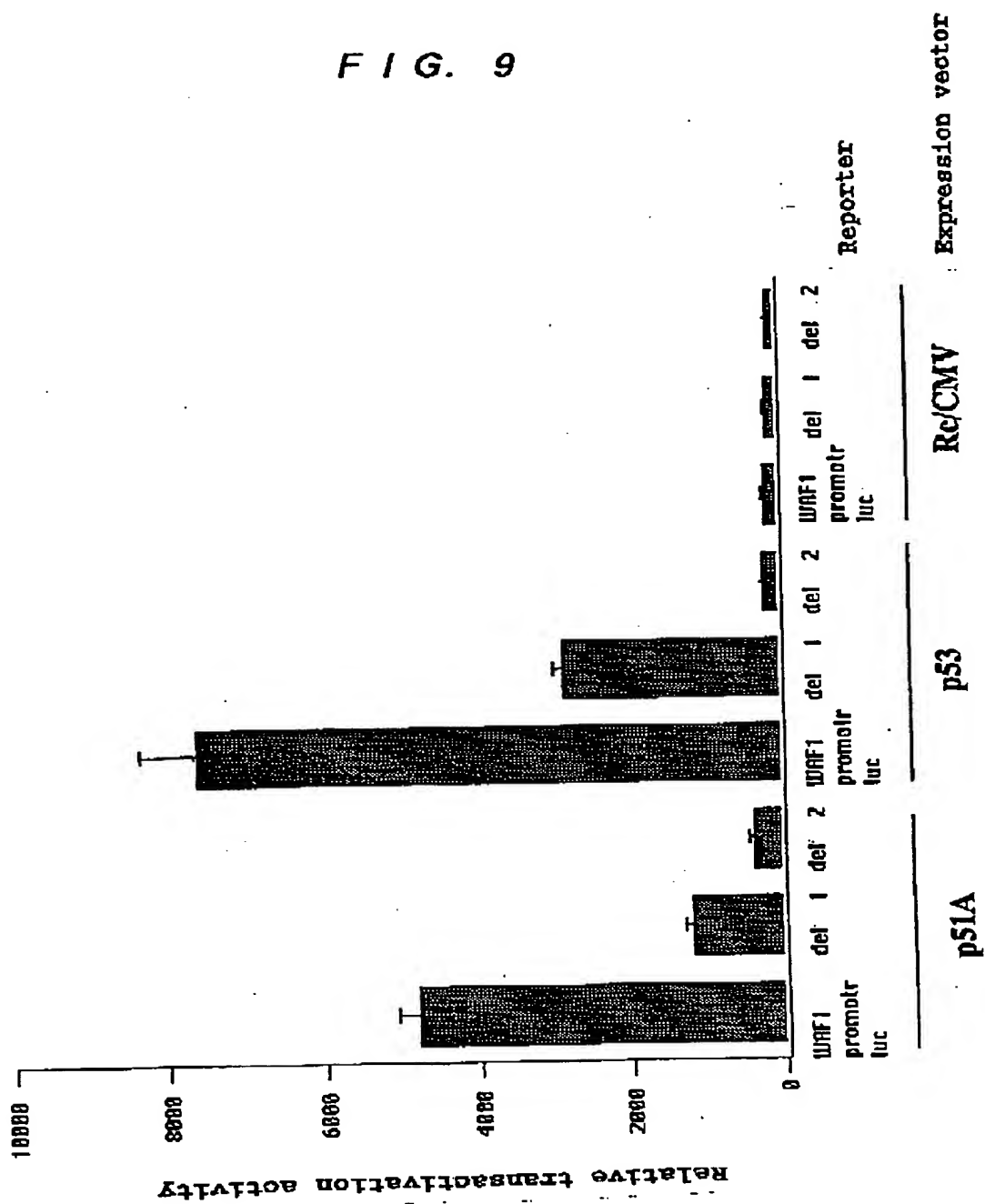


FIG. 10

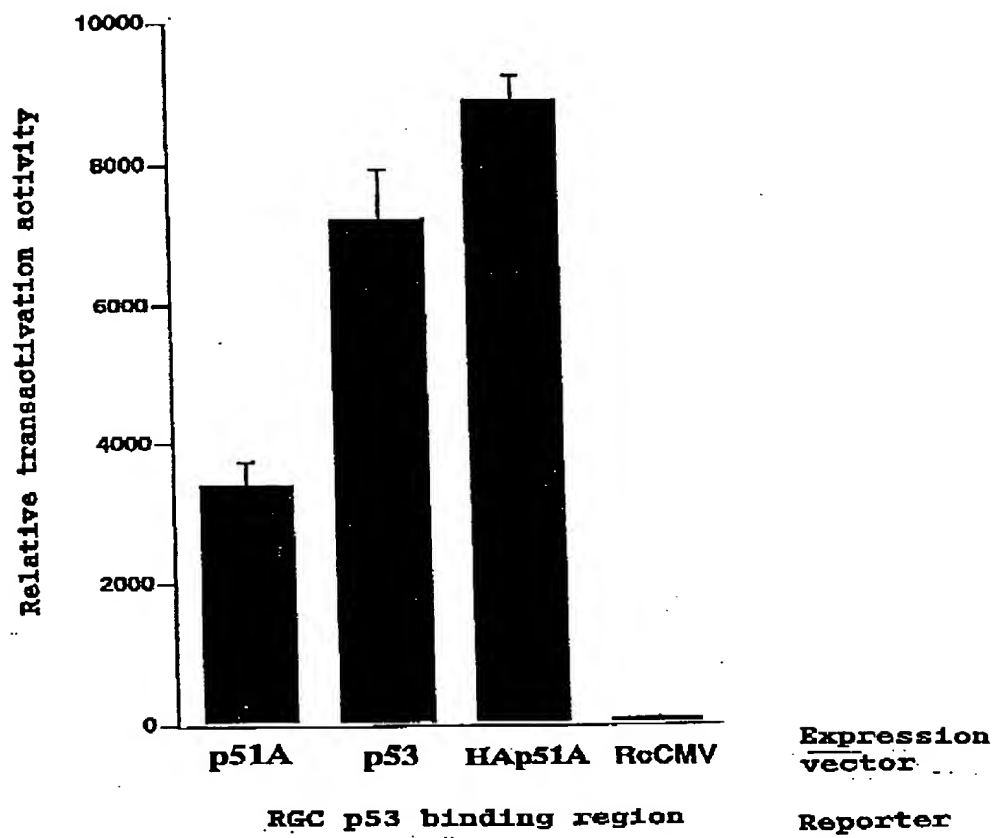


FIG. 11

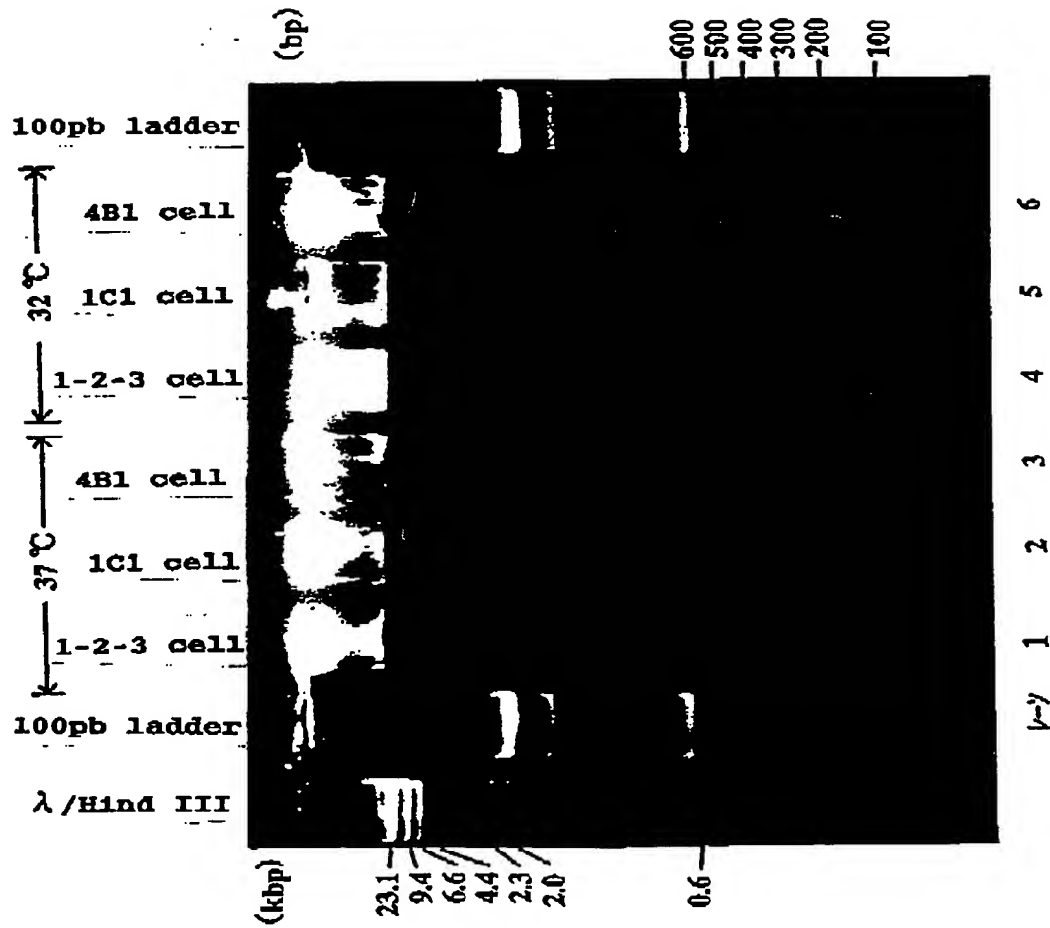


FIG. 12

mp51	1	ATGTCCAGAGCACCAGACAGCGAGTTCCTCAGCCAGAGGTCTTCCAGCATATCTGG	60
p51B	1	ATGTCCAGAGCACCAGACAGCGAGTTCCTCAGCCAGAGGTCTTCCAGCATATCTGG	60

mp51	61	GATTTTCTGGAACAGCCATATATGCTCACTACGCCCCATCGAGTTGAACTTTGTGGATGAA	120
p51B	61	GATTTTCTGGAACAGCCATATATGCTCACTACGCCCCATCGAGTTGAACTTTGTGGATGAA	120

mp51	121	CCTTCCGAAAATGCTGCAACAAACAAGATTGAGATTAGCATGGATTGTATCCGCATGCAA	180
p51B	121	CCATCAGAAAGATGCTGCGACAAACAAGATTGAGATTAGCATGGATTGTATCCGCATGCAA	180
		**	
mp51	181	GACTCAGACCTTCAGTGACCCCATGTGGCCACAGTACACGAACCTGGGGCTCCTGAACAGC	240
p51B	181	GACTCGGACCTTCAGTGACCCCATGTGGCCACAGTACACGAACCTGGGGCTCCTGAACAGC	240

mp51	241	ATGGACCAGCAGATTCAGAACGGCTCCTCGTCCACCGCCCTACAAACACAGACCACGCA	300
p51B	241	ATGGACCAGCAGATTCAGAACGGCTCCTCGTCCACCGCCCTACAAACACAGACCACGCA	300

mp51	301	CAGAAATAGCGTGACGGCGCCCTCGCCCTATGCACAGCCCAAGTCCGACCTTTGATGCCCTC	360
p51B	301	CAGAACAGCGTGACGGCGCCCTCGCCCTATGCACAGCCCAAGTCCGACCTTTGATGCCCTC	360

mp51	361	TCTCCATCCCCCTGCCATTCCTCCAAACAGATTACCCGCGCCCAACAGCTTCGATGTG	420
p51B	361	TCTCCATCCCCCTGCCATTCCTCCAAACAGATTACCCGCGCCCAACAGCTTCGATGTG	420

mp51	421	TCCTTCAGCAGTCAAGCACTGCCAAGTCAGCCACCTGGACGTATTCCACCGAAGTGAAG	480
p51B	421	TCCTTCAGCAGTCAAGCACTGCCAAGTCAGCCACCTGGACGTATTCCACCGAAGTGAAG	480

mp51	481	AAGCTGTACTGCCAAGATTGCCAAGACATGCCCCATCCAGATCAAGGTGATGACCCACCC	540
p51B	481	AAGCTGTACTGCCAAGATTGCCAAGACATGCCCCATCCAGATCAAGGTGATGACCCACCC	540
		**	
mp51	541	CCACAGGGGGCTGTTATCCGTGCCATGCCCTGTCTACAGAAAGCTGAGCATGTCCACCGAG	600
p51B	541	CCACAGGGGGCTGTTATCCGTGCCATGCCCTGTCTACAGAAAGCTGAGCATGTCCACCGAG	600
		**	
mp51	601	GTTGTGAAGCGATGCCCTAACCATGAGCTGAGCCGTGAGTTCBAATGAGGGACAGATTGCC	660
p51B	601	GTGGTGAAGCGGTGCCCTAACCATGAGCTGAGCCGTGAGTTCBAATGAGGGACAGATTGCC	660
		**	
mp51	661	CCTCCAGTCACTCTGATTGAGTAGAAGGGAAACAGCCATGCCAGTATGTAGAAGATCCCT	720
p51B	661	CCTCCAGTCACTCTGATTGAGTAGAAGGGAAACAGCCATGCCAGTATGTAGAAGATCCCT	720

mp51	721	ATCAGGGGAAGGCAGAGCGTGCTGGTCCCTTATGAGCCACCAAGGTTGGCACTGAATTC	780
p51B	721	ATCAGGGGAAGGCAGAGCGTGCTGGTCCCTTATGAGCCACCAAGGTTGGCACTGAATTC	780

FIG. 13

mp51	781	ACBACAGTCCTGTACAAATTTTCATGTGTAAACAGCAGCTGCGTCGGAGGAATGAACAGACGT	840
p51B	781	ACGACAGTCTTGTACAAATTTTCATGTGTAAACAGCAGTTGTGTGGAGGGATGAACCGCCGT	840
		** ***** ** ** ***** ***** + ***	
mp51	841	CCAATTTTAAATCATCTGTTACTCTGGAAACCBAGAGATGGGCAAGTCCTGGGCCGACGGTGC	900
p51B	841	CCAATTTTAAATCATCTGTTACTCTGGAAACCBAGAGATGGGCAAGTCCTGGGCCGACGGTGC	900

mp51	901	TTTGAGGCCCGGATCTGTGCTTGGCCAGGAAGAGACCGGAAAGGCAGATGAAGACGCCATC	960
p51B	901	TTTGAGGCCCGGATCTGTGCTTGGCCAGGAAGAGACCGGAAAGGCAGATGAAGATAGCATC	960

mp51	961	AGAAAGCAGCAAGTATCGGACAGCGCAAGGAACGGCGATGGTACGAAGCGCCCTTCCGT	1020
p51B	961	AGAAAGCAGCAAGTATCGGACAGTACAAAGAACGGTGATGGTACGAAGCGCCCGTTTCGT	1020

mp51	1021	CAGAATACACACGGAATCCAGATGACTTCCATGAGAAACGGAGATCCCGAGATGATGAG	1080
p51B	1021	CAGAACACACATGGTATCCAGATGACATCCATCAACABACGAGATCCCGAGATGATGAA	1080

mp51	1081	CTGCTGTACCTACCACTGAGAGGTGCTGAGACGTACGAGATGTTGCTGAAGATCABAGAG	1140
p51B	1081	CTGTATACTTACCACTGAGAGGTGCTGAGACGTATGAAATGCTGTTGAAGATCABAGAG	1140
		*** : *** ***** ** ***** ** ** *****	
mp51	1141	TCACTCGAGCTCATGCACTACCTCCCTCAGGACACGATCGAAACGCTACAGGCGCAGCAG	1200
p51B	1141	TCCTCGAATCATGCACTACCTCCCTCAGGACACATTCGAACGCTACAGGCAACAGCAA	1200
		** *****	
mp51	1201	CAGCAGCAGCACCAGCACCTACTTCAGAAACAGACCTCCATGCACTCTCAGTCTTTCATAT	1260
p51B	1201	CAGCAGCAGCACCAGCACCTACTTCAGAAACAGACCTCAATACAGTCTCCATCTTTCATAT	1260

mp51	1261	GGCAACAGTTCCCCACCTCTGAACAAATGAACAGCATGAACAAGCTGCCTTCCGTGAGC	1320
p51B	1261	GGTAACAGTTCCCCACCTCTGAACAAATGAACAGCATGAACAAGCTGCCTTCTGTGAGC	1320
		** *****	
mp51	1321	CAGCTTATCAACCCACAGCAGCGGAATGCCCTCACTCCCACCCATGCTGAGGGCATG	1380
p51B	1321	CAGCTTATCAACCCCTCAGCAGCGCAACGCCCTCACTCCCTACACCATTCCTGATGGCATG	1380

mp51	1381	GGAGCCAAACATTCCTATGATGGGCACCTCACATGCCAATGGCTGGAGACATGAATGGACTC	1440
p51B	1381	GGAGCCAAACATTCCTATGATGGGCACCCACATGCCAATGGCTGGAGACATGAATGGACTC	1440

mp51	1441	AGCCCTACCCAAAGCTGTCCCTCCCTCCACTCTCCATGCCCTCCACCTCCCACTGCACCCCA	1500
p51B	1441	AGCCCTACCCAAAGCTGTCCCTCCCTCCACTCTCCATGCCATGCCCTCCCACTGCACACCC	1500

mp51	1501	CCACCGCCCTACCCACAGACTGCAGCATTGTGAGTTTCTTAGCAAGCTTGGGCTGCTCA	1560
p51B	1501	CCACCTCCGTATCCCAAGATTGCAGCATTGTGAGTTTCTTAGCGAGCTTGGGCTGCTCA	1560

FIG. 14

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mp51 1561 TCATGCCTGGACTATTTACGACCCAGGGGCTGACCAACCATCTATCAGATTGAGCATTAC 1620
p51B 1561 TCATGCTCTGGACTATTTACGACCCAGGGGCTGACCAACCATCTATCAGATTGAGCATTAC 1620
*****

mp51 1621 TCCATGGATGATTTGGCAAGTCTGAAGATCCCTGAACAGTTCCGACATGCCATCTGGAG 1680
p51B 1621 TCCATGGATGATCTGGCAAGTCTGAAGATCCCTGAGCAATTTGACATGCCATCTGGAG 1680
*****

mp51 1681 GGCATCCTGGACCACAGGCGAGCTGCACGACTTCTCCTCACCTCCTCATCTCCTGAGGACC 1740
p51B 1681 GGCATCCTGGACCACCGGAGCTCCACGAAATTCCTCCCTTCTCATCTCCTGCGGACC 1740
*****

mp51 1741 CCAAGTGGTGCCTCTACCGTCACTCTCGGCTCCAGTGAGACCCGCTGAACTGTGATC 1800
p51B 1741 CCAAGCAGTGCCTCTACAGTCACTGTGGGCTCCAGTGAGACCCGGGCTGAGCGTGTATT 1800
*****

mp51 1801 GATGCCGTGCGCTTTACCCCTCCGCCAGACCATCTCTTTTCCACCCCGTGACGAGTGGAT 1860
p51B 1801 GATGCTGTGCGATTACCCCTCCGCCAGACCATCTCTTTTCCACCCCGTGAGTGGAT 1860
*****

mp51 1861 GATTTCAACTTTGACATGGATTCTGTCGCAACAAGCAGCAGCGTATCAAGAGGAAGGA 1920
p51B 1861 GACTTCAACTTTGACATGGATTGCTCGCCGCAATAGCAGCAGCGCATCAAGAGGAGGGG 1920
**

mp51 1921 GAA 1923
p51B 1921 GAG 1923
**

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FIG. 15

mp51BnAA	MSQSTQTSEFLSPEVTFQHIWDFLEQWICSVQFIELEFVDEPSENGATNKLEISMDCIKMQ	60
p51B aa	MSQSTQTNEFLSPEVTFQHIWDFLEQWICSVQFIDLEFVDEPSENGATNKLEISMDCIKMQ	60

mp51BnAA	DSDLSDFPMWFOYTNLGLLNSMDQQIQNGSSSTSPYNTDHAQNSVTAPSPFYAQPSSSTFDAL	120
p51B aa	DSDLSDFPMWFOYTNLGLLNSMDQQIQNGSSSTSPYNTDHAQNSVTAPSPFYAQPSSSTFDAL	120

mp51BnAA	SPSPAIPSNTOYPGPHSFDVSTFQSSSTAKSATWTYSTELEKLYCQIAKTCFIQIKVMTFF	180
p51B aa	SPSPAIPSNTOYPGPHSFDVSTFQSSSTAKSATWTYSTELEKLYCQIAKTCFIQIKVMTFF	180

mp51BnAA	PQGAVIRAMPVYKKAHVTEVVKRCFNEELEREFEQIAPPSSHILIRVEGNSHAQYVEDP	240
p51B aa	PQGAVIRAMPVYKKAHVTEVVKRCFNEELEREFEQIAPPSSHILIRVEGNSHAQYVEDP	240

mp51BnAA	ITGRQSVLVPIYEPFQVGTFTTLYNFMCNSSCVGGMNRRPILITVLETRDGQVLAGRC	300
p51B aa	ITGRQSVLVPIYEPFQVGTFTTLYNFMCNSSCVGGMNRRPILITVLETRDGQVLAGRC	300

mp51BnAA	FEARICACFGRDRKADEDSIRKQOVSDSARKNGDOTRPFRTQNTHECIQNTSIKRRSPDDE	360
p51B aa	FEARICACFGRDRKADEDSIRKQOVSDSARKNGDOTRPFRTQNTHECIQNTSIKRRSPDDE	360

mp51BnAA	LLYLFVVRGRETIEMLLKIKESLELMQYLPQHTIETTRQQQQQQHQLLQKQTSMSQSESY	420
p51B aa	LLYLFVVRGRETIEMLLKIKESLELMQYLPQHTIETTRQQQQQQHQLLQKQTSMSQSESY	420

mp51BnAA	GNSSPFLNKMNSMKNLPSSVSQLINPQQRNALTPPTMPEGMGANIPMMGTMPMAGDMNGL	480
p51B aa	GNSSPFLNKMNSMKNLPSSVSQLINPQQRNALTPPTMPEGMGANIPMMGTMPMAGDMNGL	480

mp51BnAA	SPTQALPFFLAMPSTSHCTPPFPYPTDCSIVSFLARLGCSSCLDYFTTQGLTTIYQIEHY	540
p51B aa	SPTQALPFFLAMPSTSHCTPPFPYPTDCSIVSFLARLGCSSCLDYFTTQGLTTIYQIEHY	540

mp51BnAA	SMODLASLKIFEQFRHAIWKGLDERQLHDFSSPPLLRTPSGASTVSVGSSETRGERVI	600
p51B aa	SMODLASLKIFEQFRHAIWKGLDERQLHDFSSPPLLRTPSGASTVSVGSSETRGERVI	600

mp51BnAA	DAVRYTLRQTISFFPRDEWDFNFDMDERRNKQORIKKEGE	641
p51B aa	DAVRYTLRQTISFFPRDEWDFNFDMDERRNKQORIKKEGE	641

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